

SEQUENCE LISTING

<110> Welcher, Andrew
Wen, Duanzhi
Kelly, Michael

<120> Interferon-Like Molecules and Uses Thereof

<130> 99,372-A

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<150> 60/169,720

<151> 1999-12-08

<160> 39

<170> PatentIn Ver. 2.0

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<212> DNA

<213> Rattus norvegicus

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<221> CDS

<222> (53)..(625)

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<222> (53)..(115)

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ctg aag tat tta tgg ctg gtg gcc ctc gtg gct cta tac att tca ccc 106
Leu Lys Tyr Leu Trp Leu Val Ala Leu Val Ala Leu Tyr Ile Ser Pro
5 10 15

atc cag tct cag aac tgt gtg tat ctg gat cat acc atc ttg gaa aac 154
Ile Gln Ser Gln Asn Cys Val Tyr Leu Asp His Thr Ile Leu Glu Asn
20 25 30

atg aaa ctt ctg agc agc atc agg acc acc ttt ccc tta aga tgt cta 202
Met Lys Leu Leu Ser Ser Ile Arg Thr Thr Phe Pro Leu Arg Cys Leu
35 40 45 50

aaa gat atc acg gat ttt gag ttt cct caa gag att ctg ctg tac gtc 250
Lys Asp Ile Thr Asp Phe Glu Phe Pro Gln Glu Ile Leu Leu Tyr Val
55 60 65

cag cat gtg aaa aag gac ata aag gca gtc acc tat cat ata tct tct 298
Gln His Val Lys Lys Asp Ile Lys Ala Val Thr Tyr His Ile Ser Ser
70 75 80

ctg gcg cta att att ttc agt ctt aaa gac tcc atc tcc ctg gcg aca 346
 Leu Ala Leu Ile Ile Phe Ser Leu Lys Asp Ser Ile Ser Leu Ala Thr
 85 90 95

gag gaa cgc ttg gaa cgt atc aga tcg gga ctt ttc aaa caa gtg cag 394
 Glu Glu Arg Leu Glu Arg Ile Arg Ser Gly Leu Phe Lys Gln Val Gln
 100 105 110

caa gct cga gag tgc atg gta gac gag gag aac aag aac acg gag gag 442
 Gln Ala Arg Glu Cys Met Val Asp Glu Glu Asn Lys Asn Thr Glu Glu
 115 120 125 130

gac agt aca tca caa cat cct cac tca gag ggc ttc aag gca gtc tac 490
 Asp Ser Thr Ser Gln His Pro His Ser Glu Gly Phe Lys Ala Val Tyr
 135 140 145

ctg gaa ttg aac aag tat ttc ttc aga atc aga aag ttc ctg gta aat 538
 Leu Glu Leu Asn Lys Tyr Phe Phe Arg Ile Arg Lys Phe Leu Val Asn
 150 155 160

aag aaa tac agt ttc tgt gcc tgg aag att gtc gtg gtg gaa ata aga 586
 Lys Lys Tyr Ser Phe Cys Ala Trp Lys Ile Val Val Val Glu Ile Arg
 165 170 175

aga tgt ttc agt ata ttt tac aaa cta ctc aac atg aat tgagaatcat 635
 Arg Cys Phe Ser Ile Phe Tyr Lys Leu Leu Asn Met Asn
 180 185 190

ccagcttcaa gcaagaactt agatagaagt tgtgactgct caaatgtccc caagaacgct 695

tgattctaag gctattgcga gtctgctgct acacacttcg gacgcaagac ttttcaagggt 755

caggggttcaa ggtagtacag tcaaaggaag tcttatgtta agcaaaagaa aaatttcagt 815

ggaaaagcta gcagaaatgt caacttgtca aaaaaacaac ttatggatta tggcattgac 875

gttactagca aaaaaaataa aacaaaaaaa aacaaaaa 913

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<212> PRT

<213> Rattus norvegicus

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 20 25 30

Glu Asn Met Lys Leu Leu Ser Ser Ile Arg Thr Thr Phe Pro Leu Arg
 35 40 45

Cys Leu Lys Asp Ile Thr Asp Phe Glu Phe Pro Gln Glu Ile Leu Leu
 50 55 60

Tyr Val Gln His Val Lys Lys Asp Ile Lys Ala Val Thr Tyr His Ile
65 70 75 80

Ser Ser Leu Ala Leu Ile Ile Phe Ser Leu Lys Asp Ser Ile Ser Leu
85 90 95

Ala Thr Glu Glu Arg Leu Glu Arg Ile Arg Ser Gly Leu Phe Lys Gln
100 105 110

Val Gln Gln Ala Arg Glu Cys Met Val Asp Glu Glu Asn Lys Asn Thr
115 120 125

Glu Glu Asp Ser Thr Ser Gln His Pro His Ser Glu Gly Phe Lys Ala
130 135 140

Val Tyr Leu Glu Leu Asn Lys Tyr Phe Phe Arg Ile Arg Lys Phe Leu
145 150 155 160

Val Asn Lys Lys Tyr Ser Phe Cys Ala Trp Lys Ile Val Val Val Glu
165 170 175

Ile Arg Arg Cys Phe Ser Ile Phe Tyr Lys Leu Leu Asn Met Asn
180 185 190

<210> 3

<211> 168

<212> PRT

<213> Rattus norvegicus

<400> 3

Cys Val Tyr Leu Asp His Thr Ile Leu Glu Asn Met Lys Leu Leu Ser
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Ser Ile Arg Thr Thr Phe Pro Leu Arg Cys Leu Lys Asp Ile Thr Asp
20 25 30

Phe Glu Phe Pro Gln Glu Ile Leu Leu Tyr Val Gln His Val Lys Lys
35 40 45

Asp Ile Lys Ala Val Thr Tyr His Ile Ser Ser Leu Ala Leu Ile Ile
50 55 60

Phe Ser Leu Lys Asp Ser Ile Ser Leu Ala Thr Glu Glu Arg Leu Glu
65 70 75 80

Arg Ile Arg Ser Gly Leu Phe Lys Gln Val Gln Gln Ala Arg Glu Cys
85 90 95

Met Val Asp Glu Glu Asn Lys Asn Thr Glu Glu Asp Ser Thr Ser Gln
100 105 110

His Pro His Ser Glu Gly Phe Lys Ala Val Tyr Leu Glu Leu Asn Lys
115 120 125

Tyr Phe Phe Arg Ile Arg Lys Phe Leu Val Asn Lys Lys Tyr Ser Phe

130

135

140

Cys Ala Trp Lys Ile Val Val Val Glu Ile Arg Arg Cys Phe Ser Ile
 145 150 155 160

Phe Tyr Lys Leu Leu Asn Met Asn
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<210> 4

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<212> DNA

<213> Homo sapiens

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<222> (575)..(1195)

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<222> (575)..(655)

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 gctatagcac caggtacaaa aaatatattt tcatgaagga tcaactccctc ttatgtaata 180
 gatttggtg agtgagtgtg tgagtgtgtg catggactca cagcttttgg ctttctgaaa 240
 taccctgcat cagtcttggt atgatgattc cttagtgtgt ggatggatca tccaggcatt 300
 taaggtaaca cgatggtaat tctttgtctc tttttcaggg aaaaaaaaaa gttatcactt 360
 ccaaagtcgg catagtcacc cgaagtaaaa aaaaaaaaaa aaaaaaaaag cctcagaggc 420
 aaaggaaagg ggccgcaacc ttggttaact gtgaaatgac gaatgagaaa actcctcctg 480
 ctgaagatat tcaggtatat aaaggcacat gaaggaaaac tcaaaacatc attgtcatat 540
 acacatcttc tggatttttt agcttgcaaa aaaa atg agc acc aaa cct gat atg 595
 Met Ser Thr Lys Pro Asp Met
 1 5

att caa aag tgt ttg tgg ctt gag atc ctt atg ggt ata ttc att gct 643
 Ile Gln Lys Cys Leu Trp Leu Glu Ile Leu Met Gly Ile Phe Ile Ala
 10 15 20

ggc acc cta tcc ctg gac tgt aac tta ctg aac gtt cac ctg aga aga 691
 Gly Thr Leu Ser Leu Asp Cys Asn Leu Leu Asn Val His Leu Arg Arg
 25 30 35

gtc acc tgg caa aat ctg aga cat ctg agt agt atg agc aat tca ttt 739
 Val Thr Trp Gln Asn Leu Arg His Leu Ser Ser Met Ser Asn Ser Phe
 40 45 50 55

cct gta gaa tgt cta cga gaa aac ata gct ttt gag ttg ccc caa gag 787
 Pro Val Glu Cys Leu Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln Glu
 60 65 70

ttt ctg caa tac acc caa cct atg aag agg gac atc aag aag gcc ttc 835
 Phe Leu Gln Tyr Thr Gln Pro Met Lys Arg Asp Ile Lys Lys Ala Phe
 75 80 85

tat gaa atg tcc cta cag gcc ttc aac atc ttc agc caa cac acc ttc 883
 Tyr Glu Met Ser Leu Gln Ala Phe Asn Ile Phe Ser Gln His Thr Phe
 90 95 100

aaa tat tgg aaa gag aga cac ctc aaa caa atc caa ata gga ctt gat 931
 Lys Tyr Trp Lys Glu Arg His Leu Lys Gln Ile Gln Ile Gly Leu Asp
 105 110 115

cag caa gca gag tac ctg aac caa tgc ttg gag gaa gac gag aat gaa 979
 Gln Gln Ala Glu Tyr Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn Glu
 120 125 130 135

aat gaa gac atg aaa gaa atg aaa gag aat gag atg aaa ccc tca gaa 1027
 Asn Glu Asp Met Lys Glu Met Lys Glu Asn Glu Met Lys Pro Ser Glu
 140 145 150

gcc agg gtc ccc cag ctg agc agc ctg gaa ctg agg aga tat ttc cac 1075
 Ala Arg Val Pro Gln Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe His
 155 160 165

agg ata gac aat ttc ctg aaa gaa aag aaa tac agt gac tgt gcc tgg 1123
 Arg Ile Asp Asn Phe Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala Trp
 170 175 180

gag att gtc cga gtg gaa atc aga aga tgt ttg tat tac ttt tac aaa 1171
 Glu Ile Val Arg Val Glu Ile Arg Arg Cys Leu Tyr Tyr Phe Tyr Lys
 185 190 195

ttt aca gct cta ttc agg agg aaa taaggatat ttttgaatt aaaattcctt 1225
 Phe Thr Ala Leu Phe Arg Arg Lys
 200 205

ttccctccga aatctctttc tccttctcct cctccatctt ctttttaagg attgtttgtgc 1285

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aatggctaga ggataggag cagagaatgt tgcaaaatgg taacatttca atgacttaac 1585

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catttctggc ccatttcctt ctcagcttgg tttgtttgaa ttgatgcttg tggaatggta 1765

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1836

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<213> Homo sapiens

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Leu Asn Val His Leu Arg Arg Val Thr Trp Gln Asn Leu Arg His Leu
35 40 45

Ser Ser Met Ser Asn Ser Phe Pro Val Glu Cys Leu Arg Glu Asn Ile
50 55 60

Ala Phe Glu Leu Pro Gln Glu Phe Leu Gln Tyr Thr Gln Pro Met Lys
65 70 75 80

Arg Asp Ile Lys Lys Ala Phe Tyr Glu Met Ser Leu Gln Ala Phe Asn
85 90 95

Ile Phe Ser Gln His Thr Phe Lys Tyr Trp Lys Glu Arg His Leu Lys
100 105 110

Gln Ile Gln Ile Gly Leu Asp Gln Gln Ala Glu Tyr Leu Asn Gln Cys
115 120 125

Leu Glu Glu Asp Glu Asn Glu Asn Glu Asp Met Lys Glu Met Lys Glu
130 135 140

Asn Glu Met Lys Pro Ser Glu Ala Arg Val Pro Gln Leu Ser Ser Leu
145 150 155 160

Glu Leu Arg Arg Tyr Phe His Arg Ile Asp Asn Phe Leu Lys Glu Lys
165 170 175

Lys Tyr Ser Asp Cys Ala Trp Glu Ile Val Arg Val Glu Ile Arg Arg
180 185 190

Cys Leu Tyr Tyr Phe Tyr Lys Phe Thr Ala Leu Phe Arg Arg Lys
195 200 205

<210> 6

<211> 178

<212> PRT

<213> Homo sapiens

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Cys Asn Leu Leu Asn Val His Leu Arg Arg Val Thr Trp Gln Asn Leu
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20 25 30

Glu Asn Ile Ala Phe Glu Leu Pro Gln Glu Phe Leu Gln Tyr Thr Gln
35 40 45

Pro Met Lys Arg Asp Ile Lys Lys Ala Phe Tyr Glu Met Ser Leu Gln
50 55 60

Ala Phe Asn Ile Phe Ser Gln His Thr Phe Lys Tyr Trp Lys Glu Arg
65 70 75 80

His Leu Lys Gln Ile Gln Ile Gly Leu Asp Gln Gln Ala Glu Tyr Leu
85 90 95

Asn Gln Cys Leu Glu Glu Asp Glu Asn Glu Asn Glu Asp Met Lys Glu
100 105 110

Met Lys Glu Asn Glu Met Lys Pro Ser Glu Ala Arg Val Pro Gln Leu
115 120 125

Ser Ser Leu Glu Leu Arg Arg Tyr Phe His Arg Ile Asp Asn Phe Leu
130 135 140

Lys Glu Lys Lys Tyr Ser Asp Cys Ala Trp Glu Ile Val Arg Val Glu
145 150 155 160

Ile Arg Arg Cys Leu Tyr Tyr Phe Tyr Lys Phe Thr Ala Leu Phe Arg
165 170 175

Arg Lys

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<212> PRT

<213> Homo sapiens

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20 25 30

Ser Ser Asn Phe Gln Cys Gln Lys Leu Leu Trp Gln Leu Asn Gly Arg
35 40 45

Leu Glu Tyr Cys Leu Lys Asp Arg Met Asn Phe Asp Ile Pro Glu Glu
50 55 60

Ile Lys Gln Leu Gln Gln Phe Gln Lys Glu Asp Ala Ala Leu Thr Ile

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70

75

80

Tyr Glu Met Leu Gln Asn Ile Phe Ala Ile Phe Arg Gln Asp Ser Ser
85 90 95

Ser Thr Gly Trp Asn Glu Thr Ile Val Glu Asn Leu Leu Ala Asn Val
100 105 110

Tyr His Gln Ile Asn His Leu Lys Thr Val Leu Glu Glu Lys Leu Glu
115 120 125

Lys Glu Asp Phe Thr Arg Gly Lys Leu Met Ser Ser Leu His Leu Lys
130 135 140

Arg Tyr Tyr Gly Arg Ile Leu His Tyr Leu Lys Ala Lys Glu Tyr Ser
145 150 155 160

His Cys Ala Trp Thr Ile Val Arg Val Glu Ile Leu Arg Asn Phe Tyr
165 170 175

Phe Ile Asn Lys Leu Thr Gly Tyr Leu Arg Asn
180 185

<210> 8

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Rat IFN-like
polypeptide cDNA insert and partial pAMG21 vector
sequence

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<221> CDS

<222> (4)..(510)

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ctg agc agc atc cgt acc acc ttt cct ctg cgt tgt ctg aaa gat atc 96
Leu Ser Ser Ile Arg Thr Thr Phe Pro Leu Arg Cys Leu Lys Asp Ile
20 25 30

acg gat ttt gag ttt cct caa gag att ctg ctg tac gtc cag cat gtg 144
Thr Asp Phe Glu Phe Pro Gln Glu Ile Leu Leu Tyr Val Gln His Val
35 40 45

aaa aag gac ata aag gca gtc acc tat cat ata tct tct ctg gcg cta 192
Lys Lys Asp Ile Lys Ala Val Thr Tyr His Ile Ser Ser Leu Ala Leu
50 55 60

att att ttc agt ctt aaa gac tcc atc tcc ctg gcg aca gag gaa cgc 240
Ile Ile Phe Ser Leu Lys Asp Ser Ile Ser Leu Ala Thr Glu Glu Arg

65

70

75

ttg gaa cgt atc aga tcg gga ctt ttc aaa caa gtg cag caa gct cga 288
 Leu Glu Arg Ile Arg Ser Gly Leu Phe Lys Gln Val Gln Gln Ala Arg
 80 85 90 95

gag tgc atg gta gac gag gag aac aag aac acg gag gag gac agt aca 336
 Glu Cys Met Val Asp Glu Glu Asn Lys Asn Thr Glu Glu Asp Ser Thr
 100 105 110

tca caa cat cct cac tca gag ggc ttc aag gca gtc tac ctg gaa ttg 384
 Ser Gln His Pro His Ser Glu Gly Phe Lys Ala Val Tyr Leu Glu Leu
 115 120 125

aac aag tat ttc ttc aga atc aga aag ttc ctg gta aat aag aaa tac 432
 Asn Lys Tyr Phe Phe Arg Ile Arg Lys Phe Leu Val Asn Lys Lys Tyr
 130 135 140

agt ttc tgt gcc tgg aag att gtc gtg gtg gaa att cgt cgt tgt ttc 480
 Ser Phe Cys Ala Trp Lys Ile Val Val Val Glu Ile Arg Arg Cys Phe
 145 150 155

agt att ttt tac aaa ctg ctg aac atg aat taatggatcc 520
 Ser Ile Phe Tyr Lys Leu Leu Asn Met Asn
 160 165

<210> 9

<211> 169

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Rat IFN-like
 polypeptide cDNA insert and partial pAMG21 vector
 sequence

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 1 5 10 15

Ser Ser Ile Arg Thr Thr Phe Pro Leu Arg Cys Leu Lys Asp Ile Thr
 20 25 30

Asp Phe Glu Phe Pro Gln Glu Ile Leu Leu Tyr Val Gln His Val Lys
 35 40 45

Lys Asp Ile Lys Ala Val Thr Tyr His Ile Ser Ser Leu Ala Leu Ile
 50 55 60

Ile Phe Ser Leu Lys Asp Ser Ile Ser Leu Ala Thr Glu Glu Arg Leu
 65 70 75 80

Glu Arg Ile Arg Ser Gly Leu Phe Lys Gln Val Gln Gln Ala Arg Glu
 85 90 95

Cys Met Val Asp Glu Glu Asn Lys Asn Thr Glu Glu Asp Ser Thr Ser

100

105

110

Gln His Pro His Ser Glu Gly Phe Lys Ala Val Tyr Leu Glu Leu Asn
 115 120 125

Lys Tyr Phe Phe Arg Ile Arg Lys Phe Leu Val Asn Lys Lys Tyr Ser
 130 135 140

Phe Cys Ala Trp Lys Ile Val Val Val Glu Ile Arg Arg Cys Phe Ser
 145 150 155 160

Ile Phe Tyr Lys Leu Leu Asn Met Asn
 165

<210> 10

<211> 520

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Rat IFN-like
 polypeptide cDNA insert and partial pAMG21 vector
 sequence

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<221> CDS

<222> (4)..(510)

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cat atg tgt gta tat ctc gat cat act atc ttg gag aat atg aaa ctt 48
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 1 5 10 15

ctg agc agc atc cgt acc acc ttt cct ctg cgt tgt ctg aaa gat atc 96
 Leu Ser Ser Ile Arg Thr Thr Phe Pro Leu Arg Cys Leu Lys Asp Ile
 20 25 30

acg gat ttt gag ttt cct caa gag att ctg ctg tac gtc cag cat gtg 144
 Thr Asp Phe Glu Phe Pro Gln Glu Ile Leu Leu Tyr Val Gln His Val
 35 40 45

aaa aag gac atc aag gca gtc acc tat cat atc tct tct ctg gcg ctg 192
 Lys Lys Asp Ile Lys Ala Val Thr Tyr His Ile Ser Ser Leu Ala Leu
 50 55 60

att att ttc agt ctt aaa gac tcc atc tcc ctg gcg aca gag gaa cgc 240
 Ile Ile Phe Ser Leu Lys Asp Ser Ile Ser Leu Ala Thr Glu Glu Arg
 65 70 75

ttg gaa cgt atc cgt tct ggt ctt ttc aaa caa gtg cag caa gct cgt 288
 Leu Glu Arg Ile Arg Ser Gly Leu Phe Lys Gln Val Gln Gln Ala Arg
 80 85 90 95

gag tgc atg gta gac gag gag aac aag aac acg gag gag gac agt aca 336
 Glu Cys Met Val Asp Glu Glu Asn Lys Asn Thr Glu Glu Asp Ser Thr
 100 105 110

tca caa cat cct cac tca gag ggc ttc aag gca gtc tac ctg gaa ttg 384
 Ser Gln His Pro His Ser Glu Gly Phe Lys Ala Val Tyr Leu Glu Leu
 115 120 125

aac aag tat ttc ttc cgt atc cgt aag ttc ctg gta aat aag aaa tac 432
 Asn Lys Tyr Phe Phe Arg Ile Arg Lys Phe Leu Val Asn Lys Lys Tyr
 130 135 140

agt ttc tgt gcc tgg aag att gtc gtg gtg gaa att cgt cgt tct ttc 480
 Ser Phe Cys Ala Trp Lys Ile Val Val Val Glu Ile Arg Arg Ser Phe
 145 150 155

agt att ttt tac aaa ctg ctg aac atg aat taatggatcc 520
 Ser Ile Phe Tyr Lys Leu Leu Asn Met Asn
 160 165

<210> 11

<211> 169

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Rat IFN-like
 polypeptide cDNA insert and partial pAMG21 vector
 sequence

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Met Cys Val Tyr Leu Asp His Thr Ile Leu Glu Asn Met Lys Leu Leu
 1 5 10 15

Ser Ser Ile Arg Thr Thr Phe Pro Leu Arg Cys Leu Lys Asp Ile Thr
 20 25 30

Asp Phe Glu Phe Pro Gln Glu Ile Leu Leu Tyr Val Gln His Val Lys
 35 40 45

Lys Asp Ile Lys Ala Val Thr Tyr His Ile Ser Ser Leu Ala Leu Ile
 50 55 60

Ile Phe Ser Leu Lys Asp Ser Ile Ser Leu Ala Thr Glu Glu Arg Leu
 65 70 75 80

Glu Arg Ile Arg Ser Gly Leu Phe Lys Gln Val Gln Gln Ala Arg Glu
 85 90 95

Cys Met Val Asp Glu Glu Asn Lys Asn Thr Glu Glu Asp Ser Thr Ser
 100 105 110

Gln His Pro His Ser Glu Gly Phe Lys Ala Val Tyr Leu Glu Leu Asn
 115 120 125

Lys Tyr Phe Phe Arg Ile Arg Lys Phe Leu Val Asn Lys Lys Tyr Ser
 130 135 140

Phe Cys Ala Trp Lys Ile Val Val Val Glu Ile Arg Arg Ser Phe Ser

145

150

155

160

Ile Phe Tyr Lys Leu Leu Asn Met Asn

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<210> 12

<211> 568

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human IFN-like
polypeptide cDNA insert and partial pAMG21 vector
sequence

<220>

<221> CDS

<222> (22)..(558)

<400> 12

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cgt gtt acc tgg caa aat ctg aga cat ctg agt agt atg agc aat tca	99	
Arg Val Thr Trp Gln Asn Leu Arg His Leu Ser Ser Met Ser Asn Ser		
	15 20 25	
ttt cct gta gaa tgt cta cga gaa aac ata gct ttt gag ttg ccc caa	147	
Phe Pro Val Glu Cys Leu Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln		
	30 35 40	
gag ttt ctg caa tac acc caa cct atg aag agg gac atc aag aag gcc	195	
Glu Phe Leu Gln Tyr Thr Gln Pro Met Lys Arg Asp Ile Lys Lys Ala		
	45 50 55	
ttc tat gaa atg tcc cta cag gcc ttc aac atc ttc agc caa cac acc	243	
Phe Tyr Glu Met Ser Leu Gln Ala Phe Asn Ile Phe Ser Gln His Thr		
	60 65 70	
ttc aaa tat tgg aaa gag aga cac ctc aaa caa atc caa ata gga ctt	291	
Phe Lys Tyr Trp Lys Glu Arg His Leu Lys Gln Ile Gln Ile Gly Leu		
	75 80 85 90	
gat cag caa gca gag tac ctg aac caa tgc ttg gag gaa gac gag aat	339	
Asp Gln Gln Ala Glu Tyr Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn		
	95 100 105	
gaa aat gaa gac atg aaa gaa atg aaa gag aat gag atg aaa ccc tca	387	
Glu Asn Glu Asp Met Lys Glu Met Lys Glu Asn Glu Met Lys Pro Ser		
	110 115 120	
gaa gcc agg gtc ccc cag ctg agc agc ctg gaa ctg agg aga tat ttc	435	
Glu Ala Arg Val Pro Gln Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe		
	125 130 135	

cac agg ata gac aat ttc ctg aaa gaa aag aaa tac agt gac tgt gcc 483
 His Arg Ile Asp Asn Phe Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala
 140 145 150

tgg gag att gtc cga gtg gaa atc cgt cgt tgc ctg tac tac ttt tac 531
 Trp Glu Ile Val Arg Val Glu Ile Arg Arg Cys Leu Tyr Tyr Phe Tyr
 155 160 165 170

aaa ttt acc gct ctg ttc cgt cgt aaa taatggatcc 568
 Lys Phe Thr Ala Leu Phe Arg Arg Lys
 175

<210> 13

<211> 179

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Rat IFN-like
 polypeptide cDNA insert and partial pAMG21 vector
 sequence

<400> 13

Met Cys Asn Leu Leu Asn Val His Leu Arg Arg Val Thr Trp Gln Asn
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Leu Arg His Leu Ser Ser Met Ser Asn Ser Phe Pro Val Glu Cys Leu
 20 25 30

Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln Glu Phe Leu Gln Tyr Thr
 35 40 45

Gln Pro Met Lys Arg Asp Ile Lys Lys Ala Phe Tyr Glu Met Ser Leu
 50 55 60

Gln Ala Phe Asn Ile Phe Ser Gln His Thr Phe Lys Tyr Trp Lys Glu
 65 70 75 80

Arg His Leu Lys Gln Ile Gln Ile Gly Leu Asp Gln Gln Ala Glu Tyr
 85 90 95

Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn Glu Asn Glu Asp Met Lys
 100 105 110

Glu Met Lys Glu Asn Glu Met Lys Pro Ser Glu Ala Arg Val Pro Gln
 115 120 125

Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe His Arg Ile Asp Asn Phe
 130 135 140

Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala Trp Glu Ile Val Arg Val
 145 150 155 160

Glu Ile Arg Arg Cys Leu Tyr Tyr Phe Tyr Lys Phe Thr Ala Leu Phe
 165 170 175

Arg Arg Lys

<210> 14

<211> 568

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human IFN-like polypeptide cDNA insert and partial pAMG21 vector sequence

<220>

<221> CDS

<222> (22)..(558)

<400> 14

tctagaaagg aggaataaca t atg tgt aac ctg ctg aac gtt cac ctg cgt 51
Met Cys Asn Leu Leu Asn Val His Leu Arg
1 5 10

cgt gtt acc tgg caa aat ctg aga cat ctg agt agt atg agc aat tca 99
Arg Val Thr Trp Gln Asn Leu Arg His Leu Ser Ser Met Ser Asn Ser
15 20 25

ttt cct gta gaa tgt cta cga gaa aac ata gct ttt gag ttg ccc caa 147
Phe Pro Val Glu Cys Leu Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln
30 35 40

gag ttc ctg caa tac acc caa cct atg aag agg gac atc aag aag gcc 195
Glu Phe Leu Gln Tyr Thr Gln Pro Met Lys Arg Asp Ile Lys Lys Ala
45 50 55

ttc tat gaa atg tcc cta cag gcc ttc aac atc ttc agc caa cac acc 243
Phe Tyr Glu Met Ser Leu Gln Ala Phe Asn Ile Phe Ser Gln His Thr
60 65 70

ttc aaa tat tgg aaa gag aga cac ctc aaa caa atc caa ata gga ctt 291
Phe Lys Tyr Trp Lys Glu Arg His Leu Lys Gln Ile Gln Ile Gly Leu
75 80 85 90

gat cag caa gca gag tac ctg aac caa tgc ttg gag gaa gac gag aat 339
Asp Gln Gln Ala Glu Tyr Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn
95 100 105

gaa aat gaa gac atg aaa gaa atg aaa gag aat gag atg aaa ccc tca 387
Glu Asn Glu Asp Met Lys Glu Met Lys Glu Asn Glu Met Lys Pro Ser
110 115 120

gaa gcc agg gtc ccc cag ctg agc agc ctg gaa ctg agg aga tat ttc 435
Glu Ala Arg Val Pro Gln Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe
125 130 135

cac agg ata gac aat ttc ctg aaa gaa aag aaa tac agt gac tgt gcc 483
His Arg Ile Asp Asn Phe Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala

140

145

150

tgg gag att gtc cga gtg gaa atc cgt cgt tct ctg tac tac ttt tac 531
 Trp Glu Ile Val Arg Val Glu Ile Arg Arg Ser Leu Tyr Tyr Phe Tyr
 155 160 165 170

aaa ttt acc gct ctg ttc cgt cgt aaa taatggatcc 568
 Lys Phe Thr Ala Leu Phe Arg Arg Lys
 175

<210> 15

<211> 179

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human IFN-like
 polypeptide cDNA insert and partial pAMG21 vector
 sequence

<400> 15

Met Cys Asn Leu Leu Asn Val His Leu Arg Arg Val Thr Trp Gln Asn
 1 5 10 15

Leu Arg His Leu Ser Ser Met Ser Asn Ser Phe Pro Val Glu Cys Leu
 20 25 30

Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln Glu Phe Leu Gln Tyr Thr
 35 40 45

Gln Pro Met Lys Arg Asp Ile Lys Lys Ala Phe Tyr Glu Met Ser Leu
 50 55 60

Gln Ala Phe Asn Ile Phe Ser Gln His Thr Phe Lys Tyr Trp Lys Glu
 65 70 75 80

Arg His Leu Lys Gln Ile Gln Ile Gly Leu Asp Gln Gln Ala Glu Tyr
 85 90 95

Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn Glu Asn Glu Asp Met Lys
 100 105 110

Glu Met Lys Glu Asn Glu Met Lys Pro Ser Glu Ala Arg Val Pro Gln
 115 120 125

Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe His Arg Ile Asp Asn Phe
 130 135 140

Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala Trp Glu Ile Val Arg Val
 145 150 155 160

Glu Ile Arg Arg Ser Leu Tyr Tyr Phe Tyr Lys Phe Thr Ala Leu Phe
 165 170 175

Arg Arg Lys

<210> 16
 <211> 556
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human IFN-like polypeptide cDNA insert and partial pAMG21 vector sequence

<220>

<221> CDS

<222> (1)..(546)

<400> 16

cat atg ctg gac tgt aac ctg ctg aac gtt cac ctg cgt cgt gtt acc	48
His Met Leu Asp Cys Asn Leu Leu Asn Val His Leu Arg Arg Val Thr	
1 5 10 15	
tgg caa aat ctg aga cat ctg agt agt atg agc aat tca ttt cct gta	96
Trp Gln Asn Leu Arg His Leu Ser Ser Met Ser Asn Ser Phe Pro Val	
20 25 30	
gaa tgt cta cga gaa aac ata gct ttt gag ttg ccc caa gag ttt ctg	144
Glu Cys Leu Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln Glu Phe Leu	
35 40 45	
caa tac acc caa cct atg aag agg gac atc aag aag gcc ttc tat gaa	192
Gln Tyr Thr Gln Pro Met Lys Arg Asp Ile Lys Lys Ala Phe Tyr Glu	
50 55 60	
atg tcc cta cag gcc ttc aac atc ttc agc caa cac acc ttc aaa tat	240
Met Ser Leu Gln Ala Phe Asn Ile Phe Ser Gln His Thr Phe Lys Tyr	
65 70 75 80	
tgg aaa gag aga cac ctc aaa caa atc caa ata gga ctt gat cag caa	288
Trp Lys Glu Arg His Leu Lys Gln Ile Gln Ile Gly Leu Asp Gln Gln	
85 90 95	
gca gag tac ctg aac caa tgc ttg gag gaa gac gag aat gaa aat gaa	336
Ala Glu Tyr Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn Glu Asn Glu	
100 105 110	
gac atg aaa gaa atg aaa gag aat gag atg aaa ccc tca gaa gcc agg	384
Asp Met Lys Glu Met Lys Glu Asn Glu Met Lys Pro Ser Glu Ala Arg	
115 120 125	
gtc ccc cag ctg agc agc ctg gaa ctg agg aga tat ttc cac agg ata	432
Val Pro Gln Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe His Arg Ile	
130 135 140	
gac aat ttc ctg aaa gaa aag aaa tac agt gac tgt gcc tgg gag att	480
Asp Asn Phe Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala Trp Glu Ile	
145 150 155 160	

gtc cga gtg gaa atc cgt cgt tgc ctg tac tac ttt tac aaa ttt acc 528
 Val Arg Val Glu Ile Arg Arg Cys Leu Tyr Tyr Phe Tyr Lys Phe Thr
 165 170 175

gct ctg ttc cgt cgt aaa taatggatcc 556
 Ala Leu Phe Arg Arg Lys
 180

<210> 17
 <211> 182
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Human IFN-like
 polypeptide cDNA insert and partial pAMG21 vector
 sequence

<400> 17
 His Met Leu Asp Cys Asn Leu Leu Asn Val His Leu Arg Arg Val Thr
 1 5 10 15
 Trp Gln Asn Leu Arg His Leu Ser Ser Met Ser Asn Ser Phe Pro Val
 20 25 30
 Glu Cys Leu Arg Glu Asn Ile Ala Phe Glu Leu Pro Gln Glu Phe Leu
 35 40 45
 Gln Tyr Thr Gln Pro Met Lys Arg Asp Ile Lys Lys Ala Phe Tyr Glu
 50 55 60
 Met Ser Leu Gln Ala Phe Asn Ile Phe Ser Gln His Thr Phe Lys Tyr
 65 70 75 80
 Trp Lys Glu Arg His Leu Lys Gln Ile Gln Ile Gly Leu Asp Gln Gln
 85 90 95
 Ala Glu Tyr Leu Asn Gln Cys Leu Glu Glu Asp Glu Asn Glu Asn Glu
 100 105 110
 Asp Met Lys Glu Met Lys Glu Asn Glu Met Lys Pro Ser Glu Ala Arg
 115 120 125
 Val Pro Gln Leu Ser Ser Leu Glu Leu Arg Arg Tyr Phe His Arg Ile
 130 135 140
 Asp Asn Phe Leu Lys Glu Lys Lys Tyr Ser Asp Cys Ala Trp Glu Ile
 145 150 155 160
 Val Arg Val Glu Ile Arg Arg Cys Leu Tyr Tyr Phe Tyr Lys Phe Thr
 165 170 175
 Ala Leu Phe Arg Arg Lys
 180

<210> 18
 <211> 11
 <212> PRT
 <213> Human immunodeficiency virus type 1

<400> 18
 Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
 1 5 10

<210> 19
 <211> 15
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Internalizing
 domain derived from HIV tat protein

<400> 19
 Gly Gly Gly Gly Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
 1 5 10 15

<210> 20
 <211> 21
 <212> DNA
 <213> Rattus norvegicus

<400> 20
 atgacactga agtatttatg g 21

<210> 21
 <211> 21
 <212> DNA
 <213> Rattus norvegicus

<400> 21
 attcatgttg agtagtttgt a 21

<210> 22
 <211> 48
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR primer
 1825-22

<400> 22
 gaataacata tgtgtgtata tctcgatcat actatcttgg agaatatg 48

<210> 23
 <211> 63

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1825-21

<400> 23

ccgcggatcc attaatcat gttcagcagt ttgtaaaaa tactgaaaca acgacgaatt 60

tcc

63

<210> 24

<211> 63

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1909-56

<400> 24

ccgcggatcc attaatcat gttcagcagt ttgtaaaaa tactgaaaga acgacgaatt 60

tcc

63

<210> 25

<211> 67

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1967-32

<400> 25

ttgatctaga aaggaggaat aacatatgtg taacctgctg aacgttcacc tgcgtcgtgt 60

tacctgg

67

<210> 26

<211> 71

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1982-14

<400> 26

ccgcggatcc attatttacg acggaacaga gcggtaaatt tgtaaaagta gtacaggcaa 60

cgacgatttc c

71

<210> 27
<211> 72
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1967-33

<400> 27
ccgcggatcc attatttacg acggaacaga gcggtaaatt tgtaaaagta gtacagagaa 60
cgacggattt cc 72

<210> 28
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
2103-87

<400> 28
aaggagcata tgctggactg taacctgctg aacgttcac 39

<210> 29
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1200-54

<400> 29
gttattgctc agcgggtggca 20

<210> 30
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1847-77

<400> 30
cccaagctta ccatgacact gaagtattta tg 32

<210> 31
<211> 33

<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1847-78

<400> 31
aaggaaaaaa gcggccgcat tcatgttgag tag

33

<210> 32
<211> 35
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1896-56

<400> 32
acgcgtcgac tcatcaattc atgttgagta gtttg

35

<210> 33
<211> 39
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1896-57

<400> 33
aaggaaaaaa gcggccgctc atcaattcat gttgagtag

39

<210> 34
<211> 31
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer
1954-45

<400> 34
acgcgtcgac ttattatttc ctctgaata g

31

<210> 35
<211> 42
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primer

1954-46

<400> 35
aaggaaaaaa gcggccgctt attatttcct cctgaataga gc 42

<210> 36
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1955-44

<400> 36
cccaagctta ccatgagcac caaacctgat atg 33

<210> 37
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1954-47

<400> 37
cccaagctta ccatgattca aaagtgtttg tggc 34

<210> 38
<211> 53
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1954-48

<400> 38
aaggaaaaaa gcggccgcgc ggcoctogat tttcctcctg aatagagctg taa 53

<210> 39
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer
1954-49

<400> 39
aaggaaaaaa gcggccgctt tcctcctgaa tagagctgta a 41